STIC Biotechnology Systems Branch

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Date Processed by STIC:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFW16

RAW SEQUENCE LISTING DATE: 04/05/2006
PATENT APPLICATION: US/10/727,358A TIME: 09:50:31

Input Set : A:\1216-1-006CIPSEQLISTREVISEDCEDTEXT.TXT

Output Set: N:\CRF4\04052006\J727358A.raw

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4 <110> APPLICANT: Kolesnick, Richard N.
 5
         King, Hong-Mei R.
 7 <120> TITLE OF INVENTION: Kinase Suppressor of Ras Inactivation
         for Therapy of Ras Mediated Tumorigenesis
11 <130> FILE REFERENCE: 1216-1-006CIP
13 <140> CURRENT APPLICATION NUMBER: 10/727,358A
14 <141> CURRENT FILING DATE: 2003-12-03
16 <150> PRIOR APPLICATION NUMBER: 60/384,228
17 <151> PRIOR FILING DATE: 2002-05-30
19 <150> PRIOR APPLICATION NUMBER: 60/460,023
                                                           Cass Not Camply
20 <151> PRIOR FILING DATE: 2003-04-03
                                                           Corrected, Diskette Needer
22 <150> PRIOR APPLICATION NUMBER: PCT/US03/16961
23 <151> PRIOR FILING DATE: 2003-05-29
25 <160> NUMBER OF SEQ ID NOS: 56
27 <170> SOFTWARE: FastSEQ for Windows Version 4.0
29 <210> SEO ID NO: 1
30 <211> LENGTH: 120
31 <212> TYPE: DNA
32 <213> ORGANISM: Homo sapiens
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36 gtgtctaacg acctcacaca gcaggagatc cggaccctag aggcaaagct ggtgaaatac 120
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40 <211> LENGTH: 41
41 <212> TYPE: PRT
42 <213> ORGANISM: Homo sapiens
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47 Thr Lys Cys Ser Val Ser Asn Asp Leu Thr Gln Glu Ile Arg Thr
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                                   25
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49 Leu Glu Ala Lys Leu Val Lys Tyr Ile
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53 <210> SEQ ID NO: 3
54 <211> LENGTH: 19
55 <212> TYPE: DNA
56 <213> ORGANISM: Homo sapiens
58 <400> SEQUENCE: 3
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59 ggcagtctgc gcgggctgc
61 <210> SEQ ID NO: 4
62 <211> LENGTH: 18
63 <212> TYPE: DNA
64 <213> ORGANISM: Homo sapiens
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75 cggaccctag aggcaaag
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78 <211> LENGTH: 19
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80 <213> ORGANISM: Artificial Sequence
82 <220> FEATURE:
83 <223> OTHER INFORMATION: antisense oligonucleotide
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86 cagcccgcgc agactgccg
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91 <213> ORGANISM: Artificial Sequence
93 <220> FEATURE:
94 <223> OTHER INFORMATION: antisense oligonucleotide
96 <400> SEQUENCE: 7
97 gaggtcgtta gacactga
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100 <211> LENGTH: 16
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102 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION: antisense oligonucleotide
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111 <211> LENGTH: 873
112 <212> TYPE: PRT
113 <213> ORGANISM: Mus musculus
115 <400> SEQUENCE: 9
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118 Gly Gly Gly Gly Ala Ala Ala Asp Gly Gly Ala Gly Ala Ala Val
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120 Ser Arg Ala Leu Gln Gln Cys Gly Gln Leu Gln Lys Leu Ile Asp Ile
           35
122 Ser Ile Gly Ser Leu Arg Gly Leu Arg Thr Lys Cys Ser Val Ser Asn
124 Asp Leu Thr Gln Gln Glu Ile Arg Thr Leu Glu Ala Lys Leu Val Lys
                        70
                                            75
126 Tyr Ile Cys Lys Gln Gln Gln Ser Lys Leu Ser Val Thr Pro Ser Asp
127
                    85
                                        90
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128 129		Thr	Ala	Glu 100	Leu	Asn	Ser	Tyr	Pro 105	Arg	Phe	Ser	Asp	Trp 110	Leu	Tyr
		Phe			Arg	Pro	Glu			Gln	Glu	Ile			Glu	Leu
	Thr	Lau	115	N1 -	T 011	T	01	120	B a.s.	03.	N1.	۲	125	T	<u>ما</u>	Mot
133		130					135			Glu		140				
	Leu 145	Arg	Arg	Trp	Gly	Ala 150	Ser	Thr	Glu	Glu	Cys 155	Ser	Arg	Leu	Gln	Gln 160
	Ala	Leu	Thr	Cys	Leu	Arg	Lys	Val	Thr	Gly	Leu	Gly	Gly	Glu	His	Lys
137					165					170					175	
138 139	Met	Asp	Ser	Gly 180	Trp	Ser	Ser	Thr	Asp 185	Ala	Arg	Asp	Ser	Ser 190	Leu	Gly
	Dro	Dro	Mat		Mot	Lou	Car	Cor		Gly	7 ~~	nl a	Glar		Sar	Thr
141	110	110	195	nap	Mec	п¢и	Ser	200	nea	GIY	Arg	ALG	205	AIG	per	2.12
142	Gln	Gly	Pro	Arg	Ser	Ile	Ser	Val	Ser	Ala	Leu	Pro	Ala	Ser	Asp	Ser
143		210					215					220				
144	Pro	Val	Pro	Gly	Leu	Ser	Glu	Gly	Leu	Ser	Asp	Ser	Cys	Ile	Pro	Leu
	225					230					235					240
146	His	Thr	Ser	Gly	Arg	Leu	Thr	Pro	Arg	Ala	Leu	His	Ser	Phe		Thr
147	_		_		245					250					255	_
148 149	Pro	Pro	Thr	Thr 260	Pro	Gln	Leu	Arg	Arg 265	His	Ala	Lys	Leu	Lys 270	Pro	Pro
_	Ara	Thr	Pro	-	Pro	Pro	Ser	Ara		Val	Phe	Gln	Leu		Pro	Ser
151			275		110		Jer	280	Lys	Val	7110	O.L.	285	200		502
152	Phe	Pro	Thr	Leu	Thr	Arg	Ser	Lys	Ser	His	Glu	Ser	Gln	Leu	Gly	Asn
153		290					295					300				
		Ile	Asp	Asp	Val	Thr	Pro	Met	Lys	Phe	Glu	Leu	Pro	His	Gly	
	305			_		310					315					320
	Pro	Gln	Leu	Val		Arg	Asp	Ile	Gly	Leu	Ser	Val	Thr	His		Phe
157	_		_	_	325	_	_			330	_		_		335	
	ser	Thr	Lys		Trp	Leu	Ser	Gln		Cys	Asn	Val	Cys		Lys	ser
159		-7.	n1	340		_	_	_	345	_			•	350		B
161	Met	116	355	GIY	vai	rys	Cys	-	HIS	Cys	Arg	Leu	365	Cys	nis	ASII
	Larg	Cve		Larg	G1	7 1 n	Dwo	360	C***	Arg	Tla	πh ⊷		T.OU	Pro	T.OIL
163	Lys	370	IIII	Llys	GIU	AId	375	MIG	Cys	Arg	116	380	FIIE	пец	FIU	Dea
	Ala		Len	Ara	Ara	ጥኮሎ		Ser	Wa l	Pro	Ser		Tle	Aan	Asn	Pro
165		5		•3	-119	390	Olu	Der	441	210	395	nop	110			400
		Asp	Ara	Ala	Ala		Pro	His	Phe	Gly		Len	Pro	Lvs	Ala	
167			5		405	Ų-u				410	****			_,_	415	
	Thr	Lvs	Lvs	Glu		Pro	Pro	Ala	Met	Asn	Len	Asp	Ser	Ser		Asn
169				420					425			F		430		
170	Pro	Ser	Ser	-	Thr	Ser	Ser	Thr		Ser	Ser	Pro	Ala		Phe	Leu
171			435				•	440					445			
172	Thr	Ser		Asn	Pro	Ser	Ser		Thr	Thr	Pro	Pro	Asn	Pro	Ser	Pro
173		450					455					460				
174	Gly	Gln	Arg	Asp	Ser	Arg	Phe	Ser	Phe	Pro	Asp	Ile	Ser	Ala	Cys	Ser
175	465					470					475			•		480
176	Gln	Ala	Ala	Pro	Leu	Ser	Ser	Thr	Ala	Asp	Ser	Thr	Arg	Leu	Asp	Asp

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490
178 Gln Pro Lys Thr Asp Val Leu Gly Val His Glu Ala Glu Ala Glu Glu
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180 Pro Glu Ala Gly Lys Ser Glu Ala Glu Asp Asp Glu Glu Asp Glu Val
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182 Asp Asp Leu Pro Ser Ser Arg Arg Pro Trp Arg Gly Pro Ile Ser Arg
                           535
184 Lys Ala Ser Gln Thr Ser Val Tyr Leu Gln Glu Trp Asp Ile Pro Phe
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                                            555
186 Glu Gln Val Glu Leu Gly Glu Pro Ile Gly Gln Gly Arg Trp Gly Arg
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                                        570
188 Val His Arg Gly Arg Trp His Gly Glu Val Ala Ile Arg Leu Leu Glu
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                                    585
190 Met Asp Gly His Asn Gln Asp His Leu Lys Leu Phe Lys Lys Glu Val
           595
                                600
192 Met Asn Tyr Arg Gln Thr Arg His Glu Asn Val Val Leu Phe Met Gly
                           615
                                                620
194 Ala Cys Met Asn Pro Pro His Leu Ala Ile Ile Thr Ser Phe Cys Lys
                       630
                                            635
196 Gly Arg Thr Leu His Ser Phe Val Arg Asp Pro Lys Thr Ser Leu Asp
                   645
                                        650
198 Ile Asn Lys Thr Arg Gln Ile Ala Gln Glu Ile Ile Lys Gly Met Gly
               660
                                    665
200 Tyr Leu His Ala Lys Gly Ile Val His Lys Asp Leu Lys Ser Lys Asn
                                680
202 Val Phe Tyr Asp Asn Gly Lys Val Val Ile Thr Asp Phe Gly Leu Phe
                            695
204 Gly Ile Ser Gly Val Val Arg Glu Glu Arg Arg Glu Asn Gln Leu Lys
                       710
                                            715
206 Leu Ser His Asp Trp Leu Cys Tyr Leu Ala Pro Glu Ile Val Arg Glu
                   725
                                        730
208 Met Ile Pro Gly Arg Asp Glu Asp Gln Leu Pro Phe Ser Lys Ala Ala
               740
                                    745
210 Asp Val Tyr Ala Phe Gly Thr Val Trp Tyr Glu Leu Gln Ala Arg Asp
212 Trp Pro Phe Lys His Gln Pro Ala Glu Ala Leu Ile Trp Gln Ile Gly
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                                                780
214 Ser Gly Glu Gly Val Arg Arg Val Leu Ala Ser Val Ser Leu Gly Lys
                       790
                                            795
216 Glu Val Gly Glu Ile Leu Ser Ala Cys Trp Ala Phe Asp Leu Gln Glu
217
                   805
                                        810
218 Arg Pro Ser Phe Ser Leu Leu Met Asp Met Leu Glu Arg Leu Pro Lys
219
               820
                                    825
220 Leu Asn Arg Arg Leu Ser His Pro Gly His Phe Trp Lys Ser Ala Asp
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222 Ile Asn Ser Ser Lys Val Met Pro Arg Phe Glu Arg Phe Gly Leu Gly
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224 Thr Leu Glu Ser Gly Asn Pro Lys Met
225 865
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228 <210> SEQ ID NO: 10
229 <211> LENGTH: 866
230 <212> TYPE: PRT
231 <213> ORGANISM: Homo sapiens
233 <400> SEQUENCE: 10
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                                   25
238 Leu Gln Lys Leu Ile Asp Ile Ser Ile Gly Ser Leu Arg Gly Leu Arg
240 Thr Lys Cys Ala Val Ser Asn Asp Leu Thr Gln Glu Ile Arg Thr
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242 Leu Glu Ala Lys Leu Val Arg Tyr Ile Cys Lys Gln Arg Gln Cys Lys
244 Leu Ser Val Ala Pro Gly Glu Arg Thr Pro Glu Leu Asn Ser Tyr Pro
246 Arg Phe Ser Asp Trp Leu Tyr Thr Phe Asn Val Arg Pro Glu Val Val
               100
                                   105
248 Gln Glu Ile Pro Arg Asp Leu Thr Leu Asp Ala Leu Leu Glu Met Asn
                               120
250 Glu Ala Lys Val Lys Glu Thr Leu Arg Arg Cys Gly Ala Ser Gly Asp
                           135
252 Glu Cys Gly Arg Leu Gln Tyr Ala Leu Thr Cys Leu Arg Lys Val Thr
                                           155
                       150
254 Gly Leu Gly Gly Glu His Lys Glu Asp Ser Ser Trp Ser Ser Leu Asp
                   165
                                       170
256 Ala Arg Arg Glu Ser Gly Ser Gly Pro Ser Thr Asp Thr Leu Ser Ala
               180
                                   185
258 Ala Ser Leu Pro Trp Pro Pro Gly Ser Ser Gln Leu Gly Arg Ala Gly
                                                   205
          195
                               200
260 Asn Ser Ala Gln Gly Pro Arg Ser Ile Ser Val Ser Ala Leu Pro Ala
                           215
                                                220
262 Ser Asp Ser Pro Thr Pro Ser Phe Ser Glu Gly Leu Ser Asp Thr Cys
                       230
                                           235
264 Ile Pro Leu His Ala Ser Gly Arg Leu Thr Pro Arg Ala Leu His Ser
                                        250
                   245
266 Phe Ile Thr Pro Pro Thr Thr Pro Gln Leu Arg Arg His Thr Lys Leu
                                   265
               260
268 Lys Pro Pro Arg Thr Pro Pro Pro Pro Ser Arg Lys Val Phe Gln Leu
           275
                               280
270 Leu Pro Ser Phe Pro Thr Leu Thr Arg Arg Lys Ser His Glu Ser Gln
                                                300
                           295
272 Leu Gly Asn Arg Ile Asp Asp Val Ser Ser Met Arg Phe Asp Leu Ser
                       310
                                            315
274 His Gly Ser Pro Gln Met Val Arg Arg Asp Ile Gly Leu Ser Val Thr
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                                       330
276 His Arg Phe Ser Thr Lys Ser Trp Leu Ser Gln Val Cys His Val Cys
277
               340
                                   345
```

727,3587

<210> 23 Dis extrain <211> 18 2113 Artificial sequence pls insent section 22207-18 22237, wheneven 22137 Lesponse is artificial on Unknown. L2237 23 atagagecea cegeatee See erron explanation on page 7.

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 04/05/2006

PATENT APPLICATION: US/10/727,358A

TIME: 09:50:32

Input Set : A:\1216-1-006CIPSEQLISTREVISEDCEDTEXT.TXT

Output Set: N:\CRF4\04052006\J727358A.raw

FRANCE EXPLANATION: 2 Use of <220> Feature (NEW RULES): Sequence(s) are missing the <220> Feature and associated headings. Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See *Federal Register, * 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seg#:23 _ VERIFICATION SUMMARY

DATE: 04/05/2006

PATENT APPLICATION: US/10/727,358A

TIME: 09:50:32

Input Set : A:\1216-1-006CIPSEQLISTREVISEDCEDTEXT.TXT

Output Set: N:\CRF4\04052006\J727358A.raw

L:604 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:23, <213>

ORGANISM: Artificial sequence

L:604 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:23, <213>

ORGANISM:Artificial sequence

L:604 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:23, Line#:604